

SEQUENCE LISTING

<110> Levy, Ilan
Shoseyov, Oded
Nussinovitch, Amos

<120> MODIFICATION OF POLYSACCHARIDE CONTAINING MATERIALS

<130> 01/22952

<160> 13

<170> PatentIn version 3.1

<210> 1

<211> 507

<212> DNA

<213> Clostridium cellulovorans

<400> 1

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caaactcaat tacaccaata atcaaaatta ctaacacatc tgacagtgtat ttaaatttaa 120

atgacgtaaa agtttagatat tattacacaa gtgatggta cacaaggacaa actttctgg 180

gtgaccatgc tgggtgcatta ttaggaaata gctatgttga taacactagc aaagtgacag 240

caaacttcgt taaagaaaaca gcaagccaa catcaaccta tgatacatat gttgaatttg 300

gatttgcaag cggacgagct actcttaaaa aaggacaatt tataactatt caaggaagaa 360

taacaaaatc agactggtca aactacactc aaacaaatga ctattcattt gatgcaagta 420

gttcaacacc agttgtaaat cccaaagtta caggatataat aggtggagct aaagtacttg 480

gtacagcacc ataggatcca gatgtac 507

<210> 2

<211> 163

<212> PRT

<213> Clostridium cellulovorans

<400> 2

Met Ala Ala Thr Ser Ser Met Ser Val Glu Phe Tyr Asn Ser Asn Lys
1 5 10 15

Ser Ala Gln Thr Asn Ser Ile Thr Pro Ile Ile Lys Ile Thr Asn Thr
20 25 30

Ser Asp Ser Asp Leu Asn Leu Asn Asp Val Lys Val Arg Tyr Tyr Tyr
35 40 45

Thr Ser Asp Gly Thr Gln Gly Gln Thr Phe Trp Cys Asp His Ala Gly
50 55 60

Ala Leu Leu Gly Asn Ser Tyr Val Asp Asn Thr Ser Lys Val Thr Ala
65 70 75 80

Asn Phe Val Lys Glu Thr Ala Ser Pro Thr Ser Thr Tyr Asp Thr Tyr
85 90 95

Val Glu Phe Gly Phe Ala Ser Gly Arg Ala Thr Leu Lys Lys Gly Gln
100 105 110

Phe Ile Thr Ile Gln Gly Arg Ile Thr Lys Ser Asp Trp Ser Asn Tyr

115 120 125

Thr Gln Thr Asn Asp Tyr Ser Phe Asp Ala Ser Ser Ser Thr Pro Val

130 135 140

Val Asn Pro Lys Val Thr Gly Tyr Ile Gly Gly Ala Lys Val Leu Gly

145 150 155 160

Thr Ala Pro

<210> 3

<211> 573

<212> DNA

<213> Clostridium cellulovorans

<400> 3

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attattacac aagtgatggc acacaaggac aaactttctg gtgtgaccat gctggtgcat 180

tatttagaaa tagctatgtt gataacacta gcaaagtgc acaaacttc gttaaagaaa 240

cagcaagccc aacatcaacc tatgatacat atgttgaatt tggatttgca agcggacgag 300

ctactcttaa aaaaggacaa ttataacta ttcaaggaag aataacaaaa tcagactggt 360

caaactacac tcaaacaaat gactattcat ttgatgcaag tagttcaaca ccagttgtaa 420
 atccaaaagt tacaggatat ataggtggag ctaaagtact tggcacagca ccaggtccag 480
 atgtaccatc ttcaataatt aatcctactt ctgcaacatt tgatcccggt accatggcta 540
 gcatgactgg tggacagcaa atgggtcgga tcc 573

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 <212> PRT
 <213> Clostridium cellulovorans

<400> 4

Met Ser Val Glu Phe Tyr Asn Ser Asn Lys Ser Ala Gln Thr Asn Ser
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Ile Thr Pro Ile Ile Lys Ile Thr Asn Thr Ser Asp Ser Asp Leu Asn
 20 25 30

Leu Asn Asp Val Lys Val Arg Tyr Tyr Thr Ser Asp Gly Thr Gln
 35 40 45

Gly Gln Thr Phe Trp Cys Asp His Ala Gly Ala Leu Leu Gly Asn Ser
 50 55 60

Tyr Val Asp Asn Thr Ser Lys Val Thr Ala Asn Phe Val Lys Glu Thr
 65 70 75 80

Ala Ser Pro Thr Ser Thr Tyr Asp Thr Tyr Val Glu Phe Gly Phe Ala
85 90 95

Ser Gly Arg Ala Thr Leu Lys Lys Gly Gln Phe Ile Thr Ile Gln Gly
100 105 110

Arg Ile Thr Lys Ser Asp Trp Ser Asn Tyr Thr Gln Thr Asn Asp Tyr
115 120 125

Ser Phe Asp Ala Ser Ser Ser Thr Pro Val Val Asn Pro Lys Val Thr
130 135 140

Gly Tyr Ile Gly Gly Ala Lys Val Leu Gly Thr Ala Pro Gly Pro Asp
145 150 155 160

Val Pro Ser Ser Ile Ile Asn Pro Thr Ser Ala Thr Phe Asp Pro Gly
165 170 175

Thr Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg Ile
180 185 190

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<213> Clostridium cellulovorans

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attattacac aagtgatggt acacaaggac aaactttctg gtgtgaccat gctggtgcat	180
tattaggaaa tagctatgtt gataacacta gcaaagtgac agcaaacttc gttaaagaaa	240
cagcaagccc aacatcaacc tatgatacat atgttgaatt tggatttgca agcggacgag	300
ctactcttaa aaaaggacaa tttataacta ttcaaggaag aataacaaaa tcagactggt	360
caaactacac tcaaacaat gactattcat ttgatgcaag tagttcaaca ccagttgtaa	420
atccaaaagt tacaggatat ataggtggag ctaaagtact tggcacagca ccaggtccag	480
atgtaccatc ttcaataatt aatcctactt ctgcaacatt tgatcccggt accatggcag	540
cgacatcatc aatgtcagtt gaattttaca actctaacaa atcagcacaa acaaactcaa	600
ttacaccaat aatcaaaatt actaacacat ctgacagtga tttaaattta aatgacgtaa	660
aagttagata ttattacaca agtgatggta cacaaggaca aactttctgg tgtgaccatg	720
ctggtgcatc attaggaaat agctatgtt ataacactag caaagtgaca gcaaacttcg	780
ttaaagaaac agcaagccc acatcaacct atgatacata tggtaattt ggatttgcaa	840
gcggacgagc tactcttaaa aaaggacaat ttataactat tcaaggaaga ataacaaaaat	900
cagactggtc aaactacact caaacaaatg actattcatt tgatgcaagt agtcaacac	960
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<211> 340

<212> PRT

<213> Clostridium cellulovorans

<400> 6

Met Ser Val Glu Phe Tyr Asn Ser Asn Lys Ser Ala Gln Thr Asn Ser

1 5 10 15

Ile Thr Pro Ile Ile Lys Ile Thr Asn Thr Ser Asp Ser Asp Leu Asn

20 25 30

Leu Asn Asp Val Lys Val Arg Tyr Tyr Thr Ser Asp Gly Thr Gln

35 40 45

Gly Gln Thr Phe Trp Cys Asp His Ala Gly Ala Leu Leu Gly Asn Ser

50 55 60

Tyr Val Asp Asn Thr Ser Lys Val Thr Ala Asn Phe Val Lys Glu Thr

65 70 75 80

Ala Ser Pro Thr Ser Thr Tyr Asp Thr Tyr Val Glu Phe Gly Phe Ala

85 90 95

Ser Gly Arg Ala Thr Leu Lys Lys Gly Gln Phe Ile Thr Ile Gln Gly

100 105 110

Arg Ile Thr Lys Ser Asp Trp Ser Asn Tyr Thr Gln Thr Asn Asp Tyr

115 120 125

Ser Phe Asp Ala Ser Ser Ser Thr Pro Val Val Asn Pro Lys Val Thr

130 135 140

Gly Tyr Ile Gly Gly Ala Lys Val Leu Gly Thr Ala Pro Gly Pro Asp

145 150 155 160

Val Pro Ser Ser Ile Ile Asn Pro Thr Ser Ala Thr Phe Asp Pro Gly

165 170 175

Thr Met Ala Ala Thr Ser Ser Met Ser Val Glu Phe Tyr Asn Ser Asn

180 185 190

Lys Ser Ala Gln Thr Asn Ser Ile Thr Pro Ile Ile Lys Ile Thr Asn

195 200 205

Thr Ser Asp Ser Asp Leu Asn Leu Asn Asp Val Lys Val Arg Tyr Tyr

210 215 220

Tyr Thr Ser Asp Gly Thr Gln Gly Gln Thr Phe Trp Cys Asp His Ala

225 230 235 240

Gly Ala Leu Leu Gly Asn Ser Tyr Val Asp Asn Thr Ser Lys Val Thr

245 250 255

Ala Asn Phe Val Lys Glu Thr Ala Ser Pro Thr Ser Thr Tyr Asp Thr

260 265 270

Tyr Val Glu Phe Gly Phe Ala Ser Gly Arg Ala Thr Leu Lys Lys Gly

275

280

285

Gln Phe Ile Thr Ile Gln Gly Arg Ile Thr Lys Ser Asp Trp Ser Asn

290

295

300

Tyr Thr Gln Thr Asn Asp Tyr Ser Phe Asp Ala Ser Ser Ser Thr Pro

305

310

315

320

Val Val Asn Pro Lys Val Thr Gly Tyr Ile Gly Gly Ala Lys Val Leu

325

330

335

Gly Thr Ala Pro

340

<210> 7

<211> 1288

<212> DNA

<213> Artificial sequence

<220>

<223> Recombinant protein sequence

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<221> misc_feature

<222> (3)..(791)

<223> Taken pRIT2T cloning vector

<220>
 <221> misc_feature
 <222> (795)..(1280)
 <223> Taken from cbpA gene

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 actctcaagc tccaaaagct gatgcgcAAC AAAATAACTT caacaaagat caacaaagcg 180
 ccttctatga aatcttgaac atgcctaact taaacgaagc gcaacgtaac ggcttcattc 240
 aaagtcttaa agacgaccca agccaaagca ctaacgtttt aggtgaagct aaaaaattaa 300
 acgaatctca agcaccgaaa gctgataaca atttcaacaa agaacaacaa aatgcttct 360
 atgaaatctt gaatatgcct aacttaaACG aagaacaacg caatggttc atccaaagct 420
 taaaagatga cccaaGCCAA agtgctaacc tattgtcaga agctaaaaag ttaaatgaat 480
 ctcaaggcacc gaaagcggat aacaattca acaaagaaca acaaaatgct ttctatgaaa 540
 tcttacattt acctaactta aacgaagaac aacgcaatgg tttcatccaa agcctaaaag 600
 atgacccaaag ccaaagcgct aaccttttag cagaagctaa aaagctaaat gatgctcaag 660
 >
 caccaaaaAGC tgacaacaaa ttcaacaaAG aacaacaaa tgctttctat gaaattttac 720
 atttacctaa cttaactgaa gaacaacgta acggcttcat ccaaaggcctt aaagacgatc 780
 cggggaaattc catggcagcg acatcatcaa tgtcagttga attttacaac tctaacaaat 840
 cagcacaaac aaactcaatt acaccaataa tcaaaattac taacacatct gacagtgatt 900

taaatttaaa tgacgtaaaa gttagatatt attacacaag tcatggtaca caaggacaaa 960
ctttctggtg tgaccatgct ggtgcattat taggaaatacg ctatgtgtat aacactagca 1020
aagtgacagc aaacttcgtt aaagaaacag caagcccaac atcaacctat gatacatatg 1080
ttgaatttgg atttgcaagc ggacgagcta ctcttaaaaa aggacaattt ataactattc 1140
aaggaagaat aacaaaatca gactggtcaa actacactca aacaaatgac tattcattt 1200
atgcaagtag ttcaacacca gttgtaaatc caaaagttac aggatataa ggtggagcta 1260
aagtacttgg tacagcacca taggatcc 1288

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<211> 426
<212> PRT
<213> Artificial sequence

<220>
<223> Recombinant protein sequence

<220>
<221> misc_feature
<222> (1)..(263)
<223> Protein A sequence, from cloning vector

<220>
<221> misc_feature
<222> (265)..(426)
<223> cbpA protein

<400> 8

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Phe Ile Gln Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Val Leu

20 25 30

Gly Glu Ala Gln Lys Leu Asn Asp Ser Gln Ala Pro Lys Ala Asp Ala

35 40 45

Gln Gln Asn Asn Phe Asn Lys Asp Gln Gln Ser Ala Phe Tyr Glu Ile

50 55 60

Leu Asn Met Pro Asn Leu Asn Glu Ala Gln Arg Asn Gly Phe Ile Gln

65 70 75 80

Ser Leu Lys Asp Asp Pro Ser Gln Ser Thr Asn Val Leu Gly Glu Ala

85 90 95

Lys Lys Leu Asn Glu Ser Gln Ala Pro Lys Ala Asp Asn Asn Phe Asn

100 105 110

Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu Asn Met Pro Asn Leu

115 120 125

Asn Glu Glu Gln Arg Asn Gly Phe Ile Gln Ser Leu Lys Asp Asp Pro

130 135 140

Ser Gln Ser Ala Asn Leu Leu Ser Glu Ala Lys Lys Leu Asn Glu Ser
145 150 155 160

Gln Ala Pro Lys Ala Asp Asn Lys Phe Asn Lys Glu Gln Gln Asn Ala
165 170 175

Phe Tyr Glu Ile Leu His Leu Pro Asn Leu Asn Glu Glu Gln Arg Asn
180 185 190

Gly Phe Ile Gln Ser Leu Lys Asp Asp Pro Ser Gln Ser Ala Asn Leu
195 200 205

Leu Ala Glu Ala Lys Lys Leu Asn Asp Ala Gln Ala Pro Lys Ala Asp
210 215 220

Asn Lys Phe Asn Lys Glu Gln Gln Asn Ala Phe Tyr Glu Ile Leu His
225 230 235 240

Leu Pro Asn Leu Thr Glu Glu Gln Arg Asn Gly Phe Ile Gln Ser Leu
245 250 255

Lys Asp Asp Pro Gly Asn Ser Met Ala Ala Thr Ser Ser Met Ser Val
260 265 270

Glu Phe Tyr Asn Ser Asn Lys Ser Ala Gln Thr Asn Ser Ile Thr Pro
275 280 285

Ile Ile Lys Ile Thr Asn Thr Ser Asp Ser Asp Leu Asn Leu Asn Asp

290 295 300

Val Lys Val Arg Tyr Tyr Thr Ser Asp Gly Thr Gln Gly Gln Thr

305 310 315 320

Phe Trp Cys Asp His Ala Gly Ala Leu Leu Gly Asn Ser Tyr Val Asp

325 330 335

Asn Thr Ser Lys Val Thr Ala Asn Phe Val Lys Glu Thr Ala Ser Pro

340 345 350

Thr Ser Thr Tyr Asp Thr Tyr Val Glu Phe Gly Phe Ala Ser Gly Arg

355 360 365

Ala Thr Leu Lys Lys Gly Gln Phe Ile Thr Ile Gln Gly Arg Ile Thr

370 375 380

Lys Ser Asp Trp Ser Asn Tyr Thr Gln Thr Asn Asp Tyr Ser Phe Asp

385 390 395 400

Ala Ser Ser Ser Thr Pro Val Val Asn Pro Lys Val Thr Gly Tyr Ile

405 410 415

Gly Gly Ala Lys Val Leu Gly Thr Ala Pro

420 425

<210> 9
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<212> DNA
<213> Artificial sequence

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<222> (68)..(624)
<223> Taken from Clostridium cellulovorans

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<221> misc_feature
<222> (652)..(981)
<223> Taken from bovine

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aactctaaca aatcagcaca aacaaactca attacaccaa taatcaaaat tactaacaca 180
tctgacagtg atttaaattt aatgacgta aaagtttagat attattacac aagtgtatgg 240
acacaaggac aaactttctg gtgtgaccat gctggtgcat tattaggaaa tagctatgtt 300
gataaacacta gcaaagtgac agcaaacttc gttaaagaaa cagcaagccc aacatcaacc 360
tatgatacat atgttgaatt tggatttgca agcggacgag ctactcttaa aaaaggacaa 420

tttataacta ttcaaggaag aataacaaaa tcagactggt caaactacac tcaaacaat	480
gactattcat ttgatgcaag tagttcaaca ccagttgtaa atccaaaagt tacaggatat	540
atagggtggag ctaaagtact tggtagcagca ccaggtccag atgtaccatc ttcaataatt	600
aatcctactt ctgcaacatt tgatcccggt accatgggtc ctcctcctgg aagcacttcc	660
gctgccagca gctccaaacta ttgcaaccag atgatgaaga gccggaacct gaccaaagat	720
cgtatgcaagc cagtgaacac ctttgtcac gagtccctgg ctgatgtcca ggcgtgtgc	780
tccccagaaaa atgtgcctg caagaatggg cagaccaatt gctaccagag ctactccacc	840
atgagcatca ccgactgccc tgagaccggc agctccaagt accccaactg tgcctacaag	900
accacccagg cgaataaaca catcattgtg gcttgtgagg gaaacccgta cgtgccagtc	960
cacttcgacg cttcagtgt a gatc	984

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 <212> PRT
 <213> Artificial sequence

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 <223> Recombinant protein sequence

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 <221> misc_feature
 <222> (30)..(208)
 <223> Taken from Clostridium cellulovorans

<220>
<221> misc_feature
<222> (226)..(326)
<223> Taken from bovine

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Ser Pro Asp Leu Gly Thr Leu Val Pro Arg Gly Ser Met Ala Ala Thr
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Ser Ser Met Ser Val Glu Phe Tyr Asn Ser Asn Lys Ser Ala Gln Thr
35 40 45

Asn Ser Ile Thr Pro Ile Ile Lys Ile Thr Asn Thr Ser Asp Ser Asp
50 55 60

Leu Asn Leu Asn Asp Val Lys Val Arg Tyr Tyr Thr Ser Asp Gly
65 70 75 80

Thr Gln Gly Gln Thr Phe Trp Cys Asp His Ala Gly Ala Leu Leu Gly
85 90 95

Asn Ser Tyr Val Asp Asn Thr Ser Lys Val Thr Ala Asn Phe Val Lys
100 105 110

Glu Thr Ala Ser Pro Thr Ser Thr Tyr Asp Thr Tyr Val Glu Phe Gly
115 120 125

Phe Ala Ser Gly Arg Ala Thr Leu Lys Lys Gly Gln Phe Ile Thr Ile
130 135 140

Gln Gly Arg Ile Thr Lys Ser Asp Trp Ser Asn Tyr Thr Gln Thr Asn
145 150 155 160

Asp Tyr Ser Phe Asp Ala Ser Ser Ser Thr Pro Val Val Asn Pro Lys
165 170 175

Val Thr Gly Tyr Ile Gly Ala Lys Val Leu Gly Thr Ala Pro Gly
180 185 190

Pro Asp Val Pro Ser Ser Ile Ile Asn Pro Thr Ser Ala Thr Phe Asp
195 200 205

Pro Gly Thr Met Gly Pro Pro Pro Gly Ser Thr Ser Ala Ala Ser Ser
210 215 220

Ser Asn Tyr Cys Asn Gln Met Met Lys Ser Arg Asn Leu Thr Lys Asp
225 230 235 240

Arg Cys Lys Pro Val Asn Thr Phe Val His Glu Ser Leu Ala Asp Val
245 250 255

Gln Ala Val Cys Ser Gln Lys Asn Val Ala Cys Lys Asn Gly Gln Thr
260 265 270

Asn Cys Tyr Gln Ser Tyr Ser Thr Met Ser Ile Thr Asp Cys Arg Glu
275 280 285

Thr Gly Ser Ser Lys Tyr Pro Asn Cys Ala Tyr Lys Thr Thr Gln Ala
290 295 300

Asn Lys His Ile Ile Val Ala Cys Glu Gly Asn Pro Tyr Val Pro Val
305 310 315 320

His Phe Asp Ala Ser Val
325

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<211> 24
<212> DNA
<213> Artificial sequence

<220>
<223> single strand DNA oligonucleotide

<400> 11
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<210> 12
<211> 18
<212> DNA
<213> Artificial sequence

<220>

<223> single strand DNA oligonucleotide

<400> 12

gggggatcct atggtgct

18

<210> 13

<211> 22

<212> DNA

<213> Artificial sequence

<220>

<223> single strand DNA oligonucleotide

<400> 13

ggggggtacc atggaacaac gc

22